Amendments to the Claims:

This listing of claims will replace all prior versions, and listings of claims in the application:

Listing of Claims:

Claims 1-48 (canceled)

Claim 49 (currently amended): A method of desorbing a macromolecular analyte 1 2 from a probe surface comprising the steps of: providing a rigid and structurally self-supporting probe that is 3 a) removably insertable into a laser desorption ionization mass spectrometer, the probe having a 4 surface for presenting the macromolecular analyte to a laser desorption ionization at least one 5 6 single energy source that emits laser energy capable of desorbing and ionizing the macromolecular analyte from the probe for analyte detection, wherein at least the surface 7 8 comprises a non-metallic-material selected from the group consisting of polystyrene, 9 polypropylene, polyethylene, polycarbonate, nylon, starch, agarose, and dextran, wherein the probe for presenting the analyte is not associated with a separate sample holder; and 10 11 b) exposing the macromolecular analyte on the probe surface to energy from a laser desorption ionization at least one single energy source, whereby the macromolecular 12 13 analyte is desorbed and ionized. Claim 50 (canceled) Claim 51 (currently amended): The method of claim 50 49 further comprising 1 2 after step (b) the steps of: 3 c) modifying the macromolecular analyte chemically or enzymatically while deposited on the probe surface; and 4 5 d) repeating step (b).

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| Claim 52 (currently amended): The method of claim 50 49 wherein the probe |
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| surface comprises an array of locations, each location having at least one macromolecular |
| analyte deposited thereon; and step (b) comprises desorbing and ionizing a first macromolecular |
| analyte from a first location in the array; |
| and wherein the method further comprises the step of (c) desorbing and ionizing a |
| second macromolecular analyte, from a second location in the array. |
| Claim 53 (currently amended): The method of claim 50 49 further comprising |
| before step (b) the step of modifying the macromolecular analyte chemically or enzymatically |
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| while deposited on the probe surface. |
| Claims 54-62 (canceled) |
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| Claim 63 (currently amended): The method of claim 50 49 wherein the |
| macromolecular analyte comprises a protein or a peptide. |
| Claim 64 (currently amended): A system for detecting a macromolecular analyte |
| comprising: |
| a removably insertable rigid and structurally self-supporting probe having a |
| surface for presenting the macromolecular analyte to a laser desorption ionization at least one |
| single energy source that emits energy capable of desorbing and ionizing the macromolecular |
| analyte from the probe, wherein at least the surface comprises a non-metallic material selected |
| from the group consisting of polystyrene, polypropylene, polyethylene, polycarbonate, nylon, |
| starch, agarose, and dextran; |
| a laser desorption ionization at least one single energy source that directs laser |
| energy to the probe surface for desorbing and ionizing the macromolecular analyte, wherein the |
| probe for presenting the analyte is not associated with a separate sample holder; and |
| a detector in communication with the probe surface that detects the desorbed |
| macromolecular analyte |

| 1 | Claim 65 (Currently amended): The system of claim 64 which is a laser |
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| 2 | desorption mass spectrometer wherein: |
| 3 | the energy source emits laser light that desorbs and ionizes the macromolecular |
| 4 | analyte to produce an ion, |
| 5 | the system further comprises means for accelerating the ion to the detector, |
| 6 | the detector detects the ion, and |
| 7 | the system further comprises means for determining the mass of the ion. |
| | Claims 66-85 (canceled) |
| 1 | Claim 86 (currently amended): A method for detecting a macromolecular analyte |
| 2 | comprising the steps of: |
| 3 | a) providing a system comprising: |
| 4 | (1) a removably insertable <u>rigid and structurally self-supporting</u> |
| 5 | probe having a surface for presenting the macromolecular analyte to a laser desorption |
| 6 | ionization at least one single energy source that emits energy capable of desorbing and ionizing |
| 7 | the macromolecular analyte from the probe, wherein at least the surface comprises a non-metallic |
| 8 | material selected from the group consisting of polystyrene, polypropylene, polyethylene, |
| 9 | polycarbonate, nylon, starch, agarose, and dextran, wherein the macromolecular analyte is |
| 10 | presented on the probe surface, wherein the probe for presenting the analyte is not associated |
| 11 | with a separate sample holder, |
| 12 | (2) <u>a laser desorption ionization</u> at least one single energy source |
| 13 | that directs <u>laser</u> energy to the probe surface for desorbing and ionizing the macromolecular |
| 14 | analyte; and |
| 15 | (3) a detector in communication with the probe surface that detects the |
| 16 | desorbed and ionized macromolecular analyte; |
| 17 | b) desorbing and ionizing at least a portion of the macromolecular analyte |
| 18 | from the surface by exposing the macromolecular analyte to energy from the laser desorption |
| 19 | ionization at least one single energy source; and |

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| 20 | c) detecting the desorbed and ionized macromolecular analyte with the |
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| 21 | detector. |
| | Cli OT () 1 1) The selection of claim of chains of the contempts of |
| 1 | Claim 87 (currently amended): The method of claim 86 wherein the system is a |
| 2 | laser desorption mass spectrometer wherein the energy source emits laser light that desorbs and |
| 3 | ionizes the macromolecular analyte to produce an ion, the detector detects the ion and the system |
| 4 | further comprises means for accelerating the ion to the detector, and the method further |
| 5 | comprises determining the mass of the ion. |
| 1 | Claim 88 (previously presented): The method of claim 87 further comprising |
| 2 | before step (b) the step of modifying the macromolecular analyte chemically or enzymatically |
| 3 | while deposited on the probe surface. |
| | |
| 1 | Claim 89 (previously presented): The method of claim 87 further comprising |
| 2 | after step (c) the steps of: |
| 3 | d) modifying the macromolecular analyte chemically or enzymatically while |
| 4 | deposited on the probe surface; and |
| 5 | e) repeating steps b) and c). |
| 1 | Claim 90 (previously presented): The method of claim 87 wherein the probe |
| 2 | surface comprises an array of locations, each location having at least one macromolecular |
| 3 | analyte deposited thereon; and step (b) comprises desorbing and ionizing a first macromolecular |
| 4 | analyte from a first location in the array; |
| 5 | and wherein the method further comprises the step of: |
| 6 | d) desorbing and ionizing a second macromolecular analyte from a second |
| 7 | location in the array; and |
| 8 | e) detecting the desorbed and ionized second macromolecular analyte with |
| 9 | the detector. |
| | nio delector. |

Claims 91-100 (canceled)

| 1 | Claim 101 (previously presented): The method of claim 87 wherein the |
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| 2 | macromolecular analyte comprises a protein or a peptide. |
| | Claims 102-104 (canceled) |
| 1 | Claim 105 (currently amended): The method of claim 50 49, wherein the |
| 2 | macromolecular analyte is a biomolecule. |
| 1 | Claim 106 (currently amended): The method of claim 50 49, wherein the |
| 2 | macromolecular analyte is a biomolecule from an undifferentiated sample. |
| 1 | Claim 107 (currently amended): The method of claim 50 49, wherein the |
| 2 | macromolecular analyte is a nucleic acid. |
| • | |
| 1 2 | Claim 108 (previously presented): The system of claim 65, wherein the macromolecular analyte is a biomolecule. |
| 2 | macromorecular analyte is a biomorecule. |
| 1 | Claim 109 (previously presented): The system of claim 65, wherein the |
| 2 | macromolecular analyte is a biomolecule from an undifferentiated sample. |
| 1 | Claim 110 (previously presented): The system of claim 65, wherein the |
| 2 | macromolecular analyte is a protein or a peptide. |
| 1 | Claim 111 (previously presented): The method of claim 87, wherein the |
| 2 | macromolecular analyte is a biomolecule. |
| | |
| 1 | Claim 112 (previously presented): The method of claim 87, wherein the |
| 2 | macromolecular analyte is a biomolecule from an undifferentiated sample. |
| 1 | Claim 113 (previously presented): The method of claim 87, wherein the |
| 2 | macromolecular analyte is a protein or a peptide. |
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Claims 114-119 (canceled)

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| 1 | Claim 120 (currently amended): The method of claim 50 49, wherein the |
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| 2 | macromolecular analyte is a carbohydrate. |
| 1 | Claim 121 (previously presented): The system of claim 65, wherein the |
| 1 | |
| 2 | macromolecular analyte is a nucleic acid. |
| 1 | Claim 122 (previously presented): The system of claim 65, wherein the |
| 2 | macromolecular analyte is a carbohydrate. |
| 1 2 | Claim 123 (previously presented): The method of claim 87, wherein the macromolecular analyte is a nucleic acid. |
| | |
| 1 | Claim 124 (previously presented): The method of claim 87, wherein the |
| 2 | macromolecular analyte is a carbohydrate. |
| | |
| 1 | Claim 125 (currently amended): The method of any of claims 49-53, 56, 57 49, |
| 2 | 51-53, 63, 105-107, 120 or 130-134 further comprising applying to the macromolecular analyte |
| 3 | associated with a matrix material for promoting desorption and ionization of the macromolecular |
| 4 | analyte on the surface. |
| 1 | China 126 (compaths are adal). The method of any of claims 64.71.75.76.92 |
| 1 | Claim 126 (currently amended): The method of any of claims 64-71, 75, 76, 82 |
| 2 | 64-65, 108-110, 121, 122 or 137-141 further comprising applying to the macromolecular analyte |
| 3 | associated with a matrix material for promoting desorption and ionization of the macromolecular |
| 4 | analyte on the surface. |
| 1 | Claim 127 (currently amended): The method of any of claims 86-91 86-90, 94, |
|) 1 | , , |
| 2 | 95, 101, 111-113, 123, 124 or 144-148 further comprising <u>applying to</u> the macromolecular |
| 3 | analyte associated with a matrix material for promoting desorption and ionization of the |
| 4 | macromolecular analyte on the surface. |

Claims 128-129 (canceled)

| 1 | Claim 130 (previously presented): The method of claim 49 wherein the non- |
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| 2 | metallic material is polystyrene. |
| 1 2 | Claim 131 (previously presented): The method of claim 49 wherein the non-metallic material is polypropylene. |
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| 1 | Claim 132 (previously presented): The method of claim 49 wherein the non- |
| 2 | metallic material is polycarbonate. |
| 1 | Claim 133 (previously presented): The method of claim 49 wherein the non- |
| 2 | metallic material is nylon. |
| 1 2 | Claim 134 (previously presented): The method of claim 49 wherein the non-metallic material is dextran. |
| | Claims 135-136 (canceled) |
| 1 2 | Claim 137 (previously presented): The system of claim 64 wherein the non-metallic material is polystyrene. |
| 1 2 | Claim 138 (previously presented): The system of claim 64 wherein the non-metallic material is polypropylene. |
| 1 2 | Claim 139 (previously presented): The system of claim 64 wherein the non-metallic material is polycarbonate. |
| 1 2 | Claim 140 (previously presented): The system of claim 64 wherein the non-metallic material is nylon. |
| 1 2 | Claim141 (previously presented): The system of claim 64 wherein the non-metallic material is dextran. |
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Claims 142-143 (canceled)

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| 1 | Claim 144 (previously presented): The method of claim 86 wherein the non- |
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| 2 | metallic material is polystyrene. |
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| 1 | Claim 145 (previously presented): The method of claim 86 wherein the non- |
| 2 | metallic material is polypropylene. |
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| 1 | Claim 146 (previously presented): The method of claim 86 wherein the non- |
| 2 | metallic material is polycarbonate. |
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| 1 | Claim 147 (previously presented): The method of claim 86 wherein the non- |
| 2 | metallic material is nylon. |
| | |
| 1 | Claim 148 (previously presented): The method of claim 86 wherein the non- |
| 2 | metallic material is dextran. |